

465 470 475 480
 Ile Gly His Val Ser Pro Glu Ala Ala Glu Gly Gly Pro Leu Ala Phe
 485 490 495
 Val Glu Asn Gly Asp His Ile Ile Val Asp Ile Glu Lys Arg Ile Leu
 500 505 510
 Asp Val Gln Val Pro Glu Glu Glu Trp Glu Lys Arg Lys Ala Asn Trp
 515 520 525
 Lys Gly Phe Glu Pro Lys Val Lys Thr Gly Tyr Leu Ala Arg Tyr Ser
 530 535 540
 Lys Leu Val Thr Ser Ala Asn Thr Gly Gly Ile Met Lys Ile
 545 550 555

<210> 39
 <211> 194
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:promoter
 sequence

<220>
 <221> -35_signal
 <222> (136)..(141)

<220>
 <221> -10_signal
 <222> (159)..(164)

<400> 39
 gctattgacg acagctatgg ttcaactgtcc accaaccaaa actgtgctca gtaccgccaa 60
 tattttctccc ttgaggggta caaagaggtg tccctagaag agatccacgc tgtgtaaaaa 120
 ttttacaaaa aggtattgac tttccctaca ggggtgtgtaa taatttaatt acaggcgggg 180
 gcaacccccgc ctgt 194

<210> 40
 <211> 163
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:promoter
 sequence

<220>
 <221> -35_signal
 <222> (113)..(118)

<220>
 <221> -10_signal
 <222> (136)..(141)

T03030"03030"

binding site

<400> 43
attcgagaaa tggagagaat ataatatg 28

<210> 44
<211> 13
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:ribosome
binding site

<400> 44
agaaaggagg tga 13

<210> 45
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:ribosome
binding site

<220>
<221> misc_feature
<222> 17-20
<223> n = a, c, g, or t

<400> 45
ttaagaaagg aggtgannnn atg 23

<210> 46
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:ribosome
binding site

<220>
<221> misc_feature
<222> 16-20
<223> n = a, c, g, or t

<400> 46
ttagaaagga ggtgannnnn atg 23

<210> 47
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<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence:ribosome
binding site